SEQUENCE LISTING

<110> Philip E. Thorpe
Rolf A. Brekken

<120> ANTIBODY CONJUGATE METHODS FOR SELECTIVELY INHIBITING VEGF

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<150> 60/131,432
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<170> PatentIn Ver. 2.0

<210> 1
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<212> DNA
<213> Homo sapiens

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<211> 498
<212> PRT
<213> Homo sapiens

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35 40 45

Glu His Asp Gly Asn Cys Arg Glu Ser Thr Thr Asp Gln Tyr Asn Thr
50 55 60

Asn Ala Leu Gln Arg Asp Ala Pro His Val Glu Pro Asp Phe Ser Ser
65 70 75 80

Gln Lys Leu Gln His Leu Glu His Val Met Glu Asn Tyr Thr Gln Trp
85 90 95

Leu Gln Lys Leu Glu Asn Tyr Ile Val Glu Asn Met Lys Ser Glu Met
100 105 110

Ala Gln Ile Gln Gln Asn Ala Val Gln Asn His Thr Ala Thr Met Leu
115 120 125

Glu Ile Gly Thr Ser Leu Leu Ser Gln Thr Ala Glu Gln Thr Arg Lys
130 135 140

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145 150 155 160

Ile Gln Leu Leu Glu Asn Ser Leu Ser Thr Tyr Lys Leu Glu Lys Gln
165 170 175

Leu Leu Gln Gln Thr Asn Glu Ile Leu Lys Ile His Glu Lys Asn Ser
180 185 190

Leu Leu Glu His Lys Ile Leu Glu Met Glu Gly Lys His Lys Glu Glu
195 200 205

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Arg Gln Thr Tyr Ile Ile Gln Glu Leu Glu Lys Gln Leu Asn Arg Ala
225 230 235 240

Thr Thr Asn Asn Ser Val Leu Gln Lys Gln Gln Leu Glu Leu Met Asp
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260 265 270

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Val Tyr Gln Ala Gly Phe Asn Lys Ser Gly Ile Tyr Thr Ile Tyr Ile
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Asn Asn Met Pro Glu Pro Lys Lys Val Phe Cys Asn Met Asp Val Asn
305 310 315 320

Gly Gly Gly Trp Thr Val Ile Gln His Arg Glu Asp Gly Ser Leu Asp
325 330 335

Phe Gln Arg Gly Trp Lys Glu Tyr Lys Met Gly Phe Gly Asn Pro Ser
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Gly Glu Tyr Trp Leu Gly Asn Glu Phe Ile Phe Ala Ile Thr Ser Gln
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Arg Gln Tyr Met Leu Arg Ile Glu Leu Met Asp Trp Glu Gly Asn Arg
370 375 380

Ala Tyr Ser Gln Tyr Asp Arg Phe His Ile Gly Asn Glu Lys Gln Asn
385 390 395 400

Tyr Arg Leu Tyr Leu Lys Gly His Thr Gly Thr Ala Gly Lys Gln Ser
405 410 415

Ser Leu Ile Leu His Gly Ala Asp Phe Ser Thr Lys Asp Ala Asp Asn
420 425 430

Asp Asn Cys Met Cys Lys Cys Ala Leu Met Leu Thr Gly Gly Trp Trp
435 440 445

Phe Asp Ala Cys Gly Pro Ser Asn Leu Asn Gly Met Phe Tyr Thr Ala
450 455 460

Gly Gln Asn His Gly Lys Leu Asn Gly Ile Lys Trp His Tyr Phe Lys
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Gly Pro Ser Tyr Ser Leu Arg Ser Thr Thr Met Met Ile Arg Pro Leu
485 490 495

Asp Phe

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<211> 2269
<212> DNA
<213> Homo sapiens

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<212> PRT
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Ala Ala Tyr Asn Asn Phe Arg Lys Ser Met Asp Ser Ile Gly Lys Lys
20 25 30

Gln Tyr Gln Val Gln His Gly Ser Cys Ser Tyr Thr Phe Leu Leu Pro
35 40 45

Glu Met Asp Asn Cys Arg Ser Ser Ser Ser Pro Tyr Val Ser Asn Ala

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     50                  55                  60

Val Gln Arg Asp Ala Pro Leu Glu Tyr Asp Asp Ser Val Gln Arg Leu
 65                  70                  75                  80

Gln Val Leu Glu Asn Ile Met Glu Asn Asn Thr Gln Trp Leu Met Lys
                85                  90                  95

Leu Glu Asn Tyr Ile Gln Asp Asn Met Lys Lys Glu Met Val Glu Ile
            100                 105                 110

Gln Gln Asn Ala Val Gln Asn Gln Thr Ala Val Met Ile Glu Ile Gly
        115                 120                 125

Thr Asn Leu Leu Asn Gln Thr Ala Glu Gln Thr Arg Lys Leu Thr Asp
    130                 135                 140

Val Glu Ala Gln Val Leu Asn Gln Thr Thr Arg Leu Glu Leu Gln Leu
145                 150                 155                 160

Leu Glu His Ser Leu Ser Thr Asn Lys Leu Glu Lys Gln Ile Leu Asp
                165                 170                 175

Gln Thr Ser Glu Ile Asn Lys Leu Gln Asp Lys Asn Ser Phe Leu Glu
            180                 185                 190

Lys Lys Val Leu Ala Met Glu Asp Lys His Ile Ile Gln Leu Gln Ser
        195                 200                 205

Ile Lys Glu Glu Lys Asp Gln Leu Gln Val Leu Val Ser Lys Gln Asn
    210                 215                 220

Ser Ile Ile Glu Glu Leu Glu Lys Lys Ile Val Thr Ala Thr Val Asn
225                 230                 235                 240

Asn Ser Val Leu Gln Lys Gln Gln His Asp Leu Met Glu Thr Val Asn
                245                 250                 255

Asn Leu Leu Thr Met Met Ser Thr Ser Asn Ser Ala Lys Asp Pro Thr
            260                 265                 270

Val Ala Lys Glu Glu Gln Ile Ser Phe Arg Asp Cys Ala Glu Val Phe
        275                 280                 285

Lys Ser Gly His Thr Thr Asn Gly Ile Tyr Thr Leu Thr Phe Pro Asn
    290                 295                 300

Ser Thr Glu Glu Ile Lys Ala Tyr Cys Asp Met Glu Ala Gly Gly Gly
305                 310                 315                 320

Gly Trp Thr Ile Ile Gln Arg Arg Glu Asp Gly Ser Val Asp Phe Gln
                325                 330                 335

Arg Thr Trp Lys Glu Tyr Lys Val Gly Phe Gly Asn Pro Ser Gly Glu
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Tyr Trp Leu Gly Asn Glu Phe Val Ser Gln Leu Thr Asn Gln Gln Arg
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        355                 360                 365

Tyr Val Leu Lys Ile His Leu Lys Asp Trp Glu Gly Asn Glu Ala Tyr
    370                 375                 380

Ser Leu Tyr Glu His Phe Tyr Leu Ser Ser Glu Glu Leu Asn Tyr Arg
385                 390                 395                 400

Ile His Leu Lys Gly Leu Thr Gly Thr Ala Gly Lys Ile Ser Ser Ile
                405                 410                 415

Ser Gln Pro Gly Asn Asp Phe Ser Thr Lys Asp Gly Asp Asn Asp Lys
            420                 425                 430

Cys Ile Cys Lys Cys Ser Gln Met Leu Thr Gly Gly Trp Trp Phe Asp
        435                 440                 445

Ala Cys Gly Pro Ser Asn Leu Asn Gly Met Tyr Tyr Pro Gln Arg Gln
    450                 455                 460

Asn Thr Asn Lys Phe Asn Gly Ile Lys Trp Tyr Tyr Trp Lys Gly Ser
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Gly Tyr Ser Leu Lys Ala Thr Thr Met Met Ile Arg Pro Ala Asp Phe
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<210> 5
<211> 495
<212> PRT
<213> Homo sapiens

<400> 5

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Ala Ala Tyr Asn Asn Phe Arg Lys Ser Met Asp Ser Ile Gly Lys Lys
            20                  25                  30

Gln Tyr Gln Val Gln His Gly Ser Cys Ser Tyr Thr Phe Leu Leu Pro
        35                  40                  45

Glu Met Asp Asn Cys Arg Ser Ser Ser Ser Pro Tyr Val Ser Asn Ala
    50                  55                  60

Val Gln Arg Asp Ala Pro Leu Glu Tyr Asp Phe Ser Ser Gln Lys Leu
65                  70                  75                  80

Gln His Leu Glu His Val Met Glu Asn Tyr Thr Gln Trp Leu Gln Lys
                85                  90                  95

Leu Glu Asn Tyr Ile Val Glu Asn Met Lys Ser Glu Met Ala Gln Ile
            100                 105                 110

Gln Gln Asn Ala Val Gln Asn His Thr Ala Thr Met Leu Glu Ile Gly
        115                 120                 125
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Thr Ser Leu Leu Ser Gln Thr Ala Glu Gln Thr Arg Lys Leu Thr Asp
    130                 135                 140

Val Glu Thr Gln Val Leu Asn Gln Thr Ser Arg Leu Glu Ile Gln Leu
145                 150                 155                 160

Leu Glu Asn Ser Leu Ser Thr Tyr Lys Leu Glu Lys Gln Leu Leu Gln
                165                 170                 175

Gln Thr Asn Glu Ile Leu Lys Ile His Glu Lys Asn Ser Leu Leu Glu
            180                 185                 190

His Lys Ile Leu Glu Met Glu Gly Lys His Lys Glu Glu Leu Asp Thr
        195                 200                 205

Leu Lys Glu Glu Lys Glu Asn Leu Gln Gly Leu Val Thr Arg Gln Thr
    210                 215                 220

Tyr Ile Ile Gln Glu Leu Glu Lys Gln Leu Asn Arg Ala Thr Thr Asn
225                 230                 235                 240

Asn Ser Val Leu Gln Lys Gln Gln Leu Glu Leu Met Asp Thr Val His
                245                 250                 255

Asn Leu Val Asn Leu Ser Thr Lys Glu Gly Val Leu Leu Lys Gly Gly
            260                 265                 270

Lys Arg Glu Glu Glu Lys Pro Phe Arg Asp Cys Ala Asp Val Tyr Gln
        275                 280                 285

Ala Gly Phe Asn Lys Ser Gly Ile Tyr Thr Ile Tyr Ile Asn Asn Met
    290                 295                 300

Pro Glu Pro Lys Lys Val Phe Cys Asn Met Asp Val Asn Gly Gly Gly
305                 310                 315                 320

Trp Thr Val Ile Gln His Arg Glu Asp Gly Ser Leu Asp Phe Gln Arg
                325                 330                 335

Gly Trp Lys Glu Tyr Lys Met Gly Phe Gly Asn Pro Ser Gly Glu Tyr
            340                 345                 350

Trp Leu Gly Asn Glu Phe Ile Phe Ala Ile Thr Ser Gln Arg Gln Tyr
        355                 360                 365

Met Leu Arg Ile Glu Leu Met Asp Trp Glu Gly Asn Arg Ala Tyr Ser
    370                 375                 380

Gln Tyr Asp Arg Phe His Ile Gly Asn Glu Lys Gln Asn Tyr Arg Leu
385                 390                 395                 400

Tyr Leu Lys Gly His Thr Gly Thr Ala Gly Lys Gln Ser Ser Leu Ile
                405                 410                 415

Leu His Gly Ala Asp Phe Ser Thr Lys Asp Ala Asp Asn Asp Asn Cys
            420                 425                 430
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Met Cys Lys Cys Ala Leu Met Leu Thr Gly Gly Trp Trp Phe Asp Ala
435 440 445

Cys Gly Pro Ser Asn Leu Asn Gly Met Phe Tyr Thr Ala Gly Gln Asn
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His Gly Lys Leu Asn Gly Ile Lys Trp His Tyr Phe Lys Gly Pro Ser
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Tyr Ser Leu Arg Ser Thr Thr Met Met Ile Arg Pro Leu Asp Phe
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<210> 6
<211> 381
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: SYNTHETIC OLIGONUCLEOTIDE

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accgtttcct ctggcggtgg c 381

<210> 7
<211> 127
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: SYNTHETIC PEPTIDE

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Gly Ala Ser Val Lys Met Ser Cys Lys Ala Ser Gly Tyr Thr Phe Thr
20 25 30

Ser Tyr Val Phe His Trp Val Lys Gln Lys Pro Gly Gln Gly Leu Glu
35 40 45

Trp Ile Gly Tyr Ile Asn Pro Tyr Asn Asp Val Thr Lys Tyr Asn Glu
50 55 60

Lys Phe Lys Gly Lys Ala Thr Leu Thr Ser Asp Lys Ser Ser Ser Thr
65 70 75 80

Ala Tyr Met Glu Leu Ser Ser Leu Thr Ser Glu Asp Ser Ala Val Tyr
85 90 95

Tyr Cys Ala Ser Tyr Tyr Gly Ser Ser Tyr Gly Tyr Tyr Ala Met Asp
100 105 110

Asp Trp Gly Gln Gly Thr Thr Val Thr Val Ser Ser Gly Gly Gly
115 120 125

<210> 8
<211> 347
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: SYNTHETIC OLIGONUCLEOTIDE

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tggtatcagc agaaaccagg gcagcctcct aaactgttga tccacggggc atccactagg 180
gaatctgggg tccctgatcg cttcacaggc agtggatctg gaaccgattt cactcttacc 240
atcagcagtg tgcaggctga agacctggca gtttattact gtcagaatga ttatagttat 300
cctctcacgt tcggtgctgg caccaagctg gaactgaaac gtctaga 347

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<211> 115
<212> PRT
<213> Artificial Sequence

<220>
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Asp Ile Gln Leu Thr Gln Ser Pro Ala Ser Leu Ser Val Ser Ala Gly
1 5 10 15

Glu Lys Val Thr Met Ser Cys Lys Ser Ser Gln Ser Leu Leu Asn Ser
20 25 30

Gly Asn Gln Lys Asn Tyr Leu Ala Trp Tyr Gln Gln Lys Pro Gly Gln
35 40 45

Pro Pro Lys Leu Leu Ile His Gly Ala Ser Thr Arg Glu Ser Gly Val
50 55 60

Pro Asp Arg Phe Thr Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr
65 70 75 80

Ile Ser Ser Val Gln Ala Glu Asp Leu Ala Val Tyr Tyr Cys Gln Asn
85 90 95

Asp Tyr Ser Tyr Pro Leu Thr Phe Gly Ala Gly Thr Lys Leu Glu Leu

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            100                 105                 110

Lys Arg Leu
        115


<210> 10
<211> 26
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:  SYNTHETIC
      PEPTIDE

<400> 10
Ala Pro Met Ala Glu Gly Gly Gly Gln Asn His His Glu Val Val Lys
  1               5                  10                  15

Phe Met Asp Val Tyr Gln Arg Ser Tyr Cys
             20                  25


<210> 11
<211> 25
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:  SYNTHETIC
      PEPTIDE

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Met Asp Val Tyr Lys Arg Ser Tyr Cys
             20                  25


<210> 12
<211> 573
<212> DNA
<213> Artificial Sequence

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      OLIGONUCLEOTIDE

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  1               5                  10                  15
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ctc cac ctg gtg gca ctg aac acc ccc ctg tct gga ggc atg cgt ggt   96
Leu His Leu Val Ala Leu Asn Thr Pro Leu Ser Gly Gly Met Arg Gly
            20                  25                  30

atc cgt gga gca gat ttc cag tgc ttc cag caa gcc cga gcc gtg ggg   144
Ile Arg Gly Ala Asp Phe Gln Cys Phe Gln Gln Ala Arg Ala Val Gly
        35                  40                  45

ctg tcg ggc acc ttc cgg gct ttc ctg tcc tct agg ctg cag gat ctc   192
Leu Ser Gly Thr Phe Arg Ala Phe Leu Ser Ser Arg Leu Gln Asp Leu
    50                  55                  60

tat agc atc gtg cgc cgt gct gac cgg ggg tct gtg ccc atc gtc aac   240
Tyr Ser Ile Val Arg Arg Ala Asp Arg Gly Ser Val Pro Ile Val Asn
65                  70                  75                  80

ctg aag gac gag gtg cta tct ccc agc tgg gac tcc ctg ttt tct ggc   288
Leu Lys Asp Glu Val Leu Ser Pro Ser Trp Asp Ser Leu Phe Ser Gly
                85                  90                  95

tcc cag ggt caa ctg caa ccc ggg gcc cgc atc ttt tct ttt gac ggc   336
Ser Gln Gly Gln Leu Gln Pro Gly Ala Arg Ile Phe Ser Phe Asp Gly
            100                 105                 110

aga gat gtc ctg aga cac cca gcc tgg ccg cag aag agc gta tgg cac   384
Arg Asp Val Leu Arg His Pro Ala Trp Pro Gln Lys Ser Val Trp His
        115                 120                 125

ggc tcg gac ccc agt ggg cgg agg ctg atg gag agt tac tgt gag aca   432
Gly Ser Asp Pro Ser Gly Arg Arg Leu Met Glu Ser Tyr Cys Glu Thr
    130                 135                 140

tgg cga act gaa act act ggg gct aca ggt cag gcc tcc tcc ctg ctg   480
Trp Arg Thr Glu Thr Thr Gly Ala Thr Gly Gln Ala Ser Ser Leu Leu
145                 150                 155                 160

tca ggc agg ctc ctg gaa cag aaa gct gcg agc tgc cac aac agc tac   528
Ser Gly Arg Leu Leu Glu Gln Lys Ala Ala Ser Cys His Asn Ser Tyr
                165                 170                 175

atc gtc ctg tgc att gag aat agc ttc atg acc tct ttc tcc aaa       573
Ile Val Leu Cys Ile Glu Asn Ser Phe Met Thr Ser Phe Ser Lys
            180                 185                 190
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<210> 13
<211> 191
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: SYNTHETIC PEPTIDE

<400> 13

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Met His His His His His His His Thr His Gln Asp Phe Gln Pro Val
  1               5                  10                  15
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Leu His Leu Val Ala Leu Asn Thr Pro Leu Ser Gly Gly Met Arg Gly
            20                  25                  30

Ile Arg Gly Ala Asp Phe Gln Cys Phe Gln Gln Ala Arg Ala Val Gly
        35                  40                  45

Leu Ser Gly Thr Phe Arg Ala Phe Leu Ser Ser Arg Leu Gln Asp Leu
    50                  55                  60

Tyr Ser Ile Val Arg Arg Ala Asp Arg Gly Ser Val Pro Ile Val Asn
65                  70                  75                  80

Leu Lys Asp Glu Val Leu Ser Pro Ser Trp Asp Ser Leu Phe Ser Gly
                85                  90                  95

Ser Gln Gly Gln Leu Gln Pro Gly Ala Arg Ile Phe Ser Phe Asp Gly
            100                 105                 110

Arg Asp Val Leu Arg His Pro Ala Trp Pro Gln Lys Ser Val Trp His
        115                 120                 125

Gly Ser Asp Pro Ser Gly Arg Arg Leu Met Glu Ser Tyr Cys Glu Thr
    130                 135                 140

Trp Arg Thr Glu Thr Thr Gly Ala Thr Gly Gln Ala Ser Ser Leu Leu
145                 150                 155                 160

Ser Gly Arg Leu Leu Glu Gln Lys Ala Ala Ser Cys His Asn Ser Tyr
                165                 170                 175

Ile Val Leu Cys Ile Glu Asn Ser Phe Met Thr Ser Phe Ser Lys
            180                 185                 190
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<210> 14
<211> 182
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: SYNTHETIC PEPTIDE

<400> 14

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His Ser His Arg Asp Phe Gln Pro Val Leu His Leu Val Ala Leu Asn
  1               5                  10                  15

Ser Pro Leu Ser Gly Gly Met Arg Gly Ile Arg Gly Ala Asp Phe Gln
            20                  25                  30

Cys Phe Gln Gln Ala Arg Ala Val Gly Leu Ala Gly Thr Phe Arg Ala
        35                  40                  45

Phe Leu Ser Ser Arg Leu Gln Asp Leu Tyr Ser Ile Val Arg Arg Ala
    50                  55                  60
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Asp Arg Ala Ala Val Pro Ile Val Asn Leu Lys Asp Glu Leu Leu Phe
 65                  70                  75                  80

Pro Ser Trp Glu Ala Leu Phe Ser Gly Ser Glu Gly Pro Leu Lys Pro
                85                  90                  95

Gly Ala Arg Ile Phe Ser Phe Asp Gly Lys Asp Val Leu Arg His Pro
            100                 105                 110

Thr Trp Pro Gln Lys Ser Val Trp His Gly Ser Asp Pro Asn Gly Arg
        115                 120                 125

Arg Leu Thr Glu Ser Tyr Cys Glu Thr Trp Arg Thr Glu Ala Pro Ser
    130                 135                 140

Ala Thr Gly Gln Ala Ser Ser Leu Leu Gly Gly Arg Leu Leu Gly Gln
145                 150                 155                 160

Ser Ala Ala Ser Cys His His Ala Tyr Ile Val Leu Cys Ile Glu Asn
                165                 170                 175

Ser Phe Met Thr Ala Ser
            180
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<210> 15
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<220>
<223> Description of Artificial Sequence: SYNTHETIC PEPTIDE

<400> 15
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1 5

<210> 16
<211> 8
<212> PRT
<213> Artificial Sequence

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<223> Description of Artificial Sequence: SYNTHETIC PEPTIDE

<400> 16
Pro Arg Phe Arg Ile Ile Gly Gly
1 5

<210> 17
<211> 9
<212> PRT
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<220>
<223> Description of Artificial Sequence: SYNTHETIC PEPTIDE

<400> 17
Ser Ser Arg His Arg Arg Ala Leu Asp
1 5

<210> 18
<211> 14
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: SYNTHETIC PEPTIDE

<400> 18
Arg Lys Ser Ser Ile Ile Ile Arg Met Arg Asp Val Val Leu
1 5 10

<210> 19
<211> 15
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: SYNTHETIC PEPTIDE

<400> 19
Ser Ser Ser Phe Asp Lys Gly Lys Tyr Lys Lys Gly Asp Asp Ala
1 5 10 15

<210> 20
<211> 15
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: SYNTHETIC PEPTIDE

<400> 20
Ser Ser Ser Phe Asp Lys Gly Lys Tyr Lys Arg Gly Asp Asp Ala
1 5 10 15

<210> 21
<211> 4
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: SYNTHETIC
PEPTIDE

<400> 21
Ile Glu Gly Arg
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<210> 22
<211> 4
<212> PRT
<213> Artificial Sequence

<220>
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PEPTIDE

<400> 22
Ile Asp Gly Arg
1

<210> 23
<211> 7
<212> PRT
<213> Artificial Sequence

<220>
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PEPTIDE

<400> 23
Gly Gly Ser Ile Asp Gly Arg
1 5

<210> 24
<211> 6
<212> PRT
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<220>
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PEPTIDE

<400> 24
Pro Leu Gly Leu Trp Ala
1 5

<210> 25
<211> 8
<212> PRT
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<220>

<223> Description of Artificial Sequence: SYNTHETIC
PEPTIDE

<400> 25
Gly Pro Gln Gly Ile Ala Gly Gln
1 5

<210> 26
<211> 8
<212> PRT
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<220>
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PEPTIDE

<400> 26
Gly Pro Gln Gly Leu Leu Gly Ala
1 5

<210> 27
<211> 5
<212> PRT
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<220>
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PEPTIDE

<400> 27
Gly Ile Ala Gly Gln
1 5

<210> 28
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
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PEPTIDE

<400> 28
Gly Pro Leu Gly Ile Ala Gly Ile
1 5

<210> 29
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
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PEPTIDE

<400> 29
Gly Pro Glu Gly Leu Arg Val Gly
1 5

<210> 30
<211> 8
<212> PRT
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<220>
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PEPTIDE

<400> 30
Tyr Gly Ala Gly Leu Gly Val Val
1 5

<210> 31
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
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PEPTIDE

<400> 31
Ala Gly Leu Gly Val Val Glu Arg
1 5

<210> 32
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
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PEPTIDE

<400> 32
Ala Gly Leu Gly Ile Ser Ser Thr
1 5

<210> 33
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
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PEPTIDE

<400> 33
Glu Pro Gln Ala Leu Ala Met Ser
1 5

<210> 34
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
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<400> 34
Gln Ala Leu Ala Met Ser Ala Ile
1 5

<210> 35
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
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<400> 35
Ala Ala Tyr His Leu Val Ser Gln
1 5

<210> 36
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
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<400> 36
Met Asp Ala Phe Leu Glu Ser Ser
1 5

<210> 37
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: SYNTHETIC PEPTIDE

<400> 37
Glu Ser Leu Pro Val Val Ala Val
1 5

<210> 38
<211> 8
<212> PRT
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<220>
<223> Description of Artificial Sequence: SYNTHETIC PEPTIDE

<400> 38
Ser Ala Pro Ala Val Glu Ser Glu
1 5

<210> 39
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
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<400> 39
Asp Val Ala Gln Phe Val Leu Thr
1 5

<210> 40
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
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<400> 40
Val Ala Gln Phe Val Leu Thr Glu
1 5

<210> 41
<211> 8
<212> PRT
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<220>
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<400> 41

Ala Gln Phe Val Leu Thr Glu Gly
1 5

<210> 42
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: SYNTHETIC PEPTIDE

<400> 42
Pro Val Gln Pro Ile Gly Pro Gln
1 5

<210> 43
<211> 31
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: SYNTHETIC OLIGONUCLEOTIDE

<400> 43
agaccatggg tcatactcat caggactttc a 31

<210> 44
<211> 29
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: SYNTHETIC OLIGONUCLEOTIDE

<400> 44
ctaccatggc tatttggaga aagaggtca 29